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Anti-collision methods for global SAW RFID **tag** systems

C Hartmann, P Hartmann, P Brown, J ... - 2004 IEEE ... , 2004 - ieeexplore.ieee.org

... Abstract - The Global SAW **Tag** [1] is projected to be a large-volume SAW device application.

The modulation method and overall **tag data structure** are reviewed for a family of proposed radio frequency identification (RFID) devices. ... II. Overall **Tag Data Structure** ...

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20, A genetic algorithm for the job shop problem – Croce, Tadei, et al. ...

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Compiler-decided dynamic memory allocation for scratch-pad based embedded systems

S Udayakumaran, R Barua - Proceedings of the 2003 international ... , 2003 - portal.acm.org

... The **tag**, **data** and valid bits are all managed by compiler-inserted software at each program **data** access ... Figure 1(b) shows the **Data** Program Relationship Graph (DPRG) for the program in figure 1(a). The DPRG is a new **data structure** we introduce that helps in the identification ...

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Garbage collection in a large LISP system

DA Moon - Proceedings of the 1984 ACM Symposium on LISP and ... , 1984 - portal.acm.org

... If the **tag** field says that the word contains a meaningful address, then the address is translated into a space by table lookup. ... If the barrier were implemented in software, it would be very expensive, since every primitive **data structure** accessing operation, such as CAR, AREF, or ...

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Compiler support for machine-independent parallel programming in Fortran D

S Hiranandani, K Kennedy, CW Tseng - Languages, Compilers, and Run ... , 1992 - Citeseer

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The GENIA corpus: An annotated research abstract corpus in molecular biology domain

T Ohta, Y Tateisi, JD Kim - ... of the second international conference on ... , 2002 - portal.acm.org

... crete **data** model (ontology) of the biological domain on which the **tag** sets are based. In defining the ontology, we required that the classification must be mutually exclusive. For example, the substances can be classified by their biological role or by their chemical **structure**, ...

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[HTML] In silico identification and expression of SLC 30 family genes: An expressed sequence tag data mining strategy for the characterization of zinc transporters' ...

M Seve, F Chimienti, S Devergnas, A Favier - BMC genomics, 2004 - biomedcentral.com

... Despite an overall shared topological **structure**, the similarity between the subfamilies is relatively low. ... The in silico characterization of SLC30 tissue expression pattern was performed by an expressed sequence **tag** (EST) **data** mining strategy. ...

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ML Fredman, J Komlós, E Szemerédi - Journal of the ACM (JACM), 1984 - portal.acm.org

... (These **tag** bits can be packed into $O(n/\log m) = o(n)$ words.) This approach requires an auxiliary **data structure** to determine whether a block W_j is nonempty, and to find the cell and **tag** bit associated with W_{\sim} when W_j is nonempty. ...

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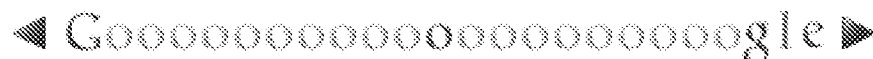
IEPAD: information extraction based on pattern discovery

CH Chang, SC Lui - ... of the 10th international conference on World ..., 2001 - portal.acm.org

... one serious problem in this one-**tag** separator approach arises when the separator **tag** is used ... target pattern may represent a desired **data** record, and all of the desired **data** records form a ... observation that desired information in a Web page is often placed in a **structure** having a ...

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